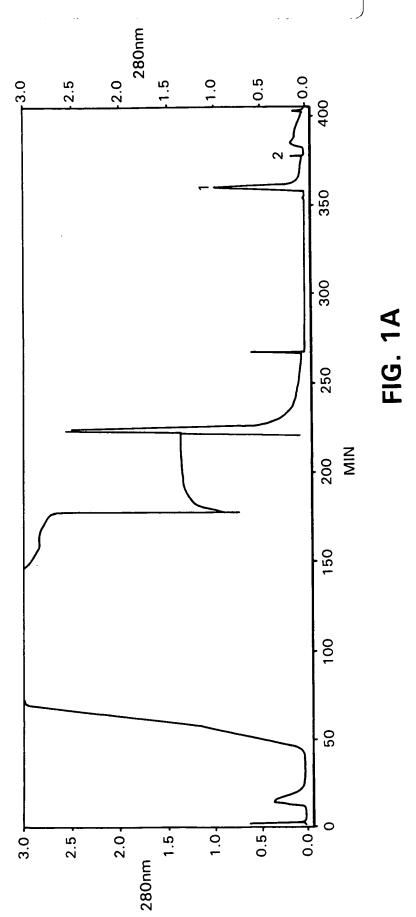


Appln No.: 10/808,041 Page 1 of Applicant(s): Roman M. Chicz et al.
PROFILING AND CATALOGING EXPRESSED PROTEIN



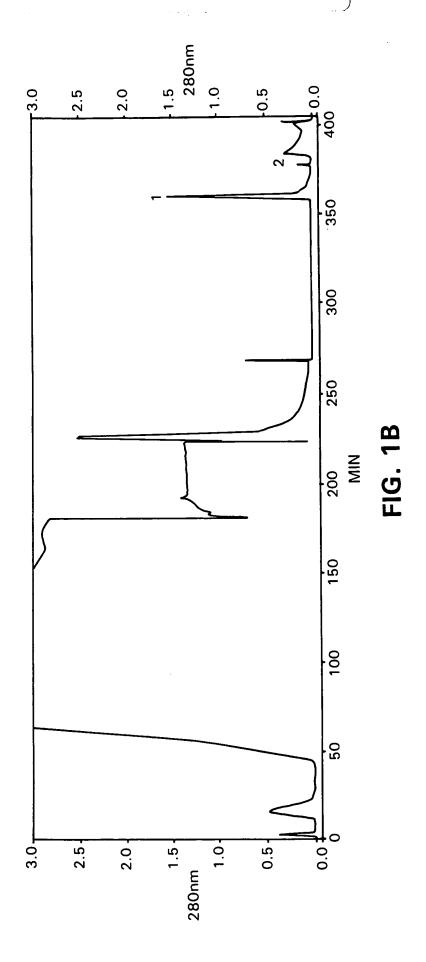
Page 1 of 10



Page 2 of 10

Applin No.: 10/808,041 Page 2 of Applicant(s): Roman M. Chicz et al. PROFILING AND CATALOGING EXPRESSED PROTEIN

TAGS



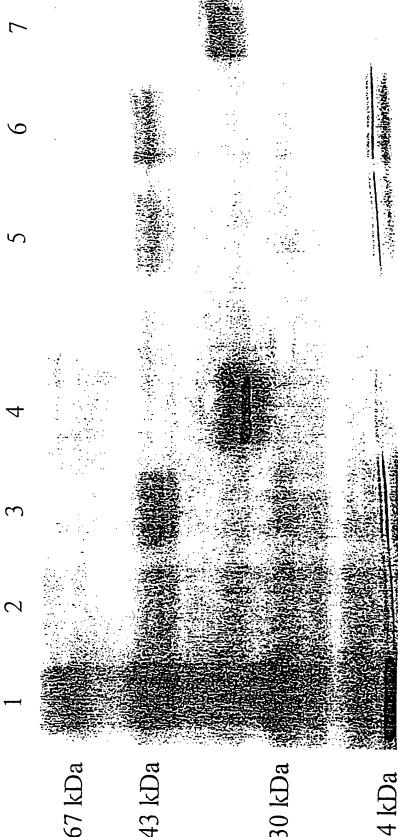
Page 3 of 10

Appln No.: 10/808,041 Page 3 of Applicant(s): Roman M. Chicz et al. PROFILING AND CATALOGING EXPRESSED PROTEIN

TAGS

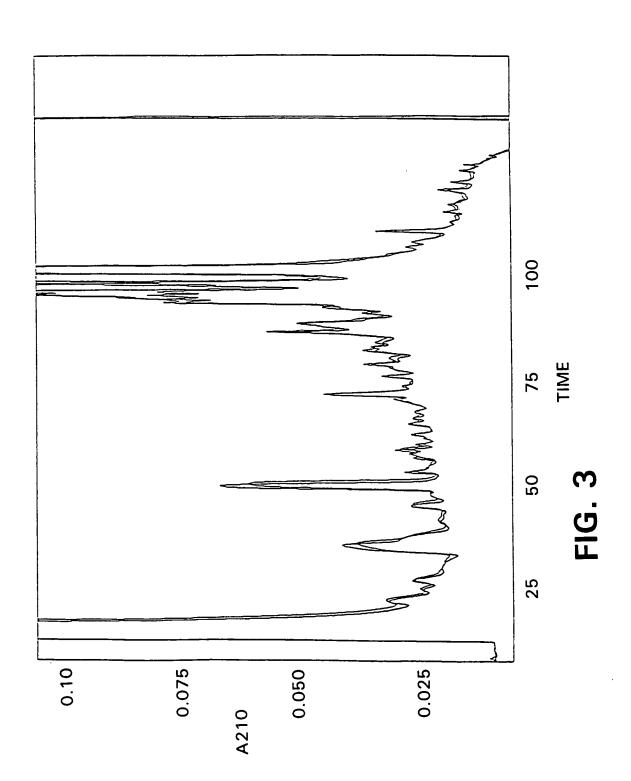
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Page 4 of 10

Appln No.: 10/808,041 Page 4 of Applicant(s): Roman M. Chicz et al. PROFILING AND CATALOGING EXPRESSED PROTEIN TAGS



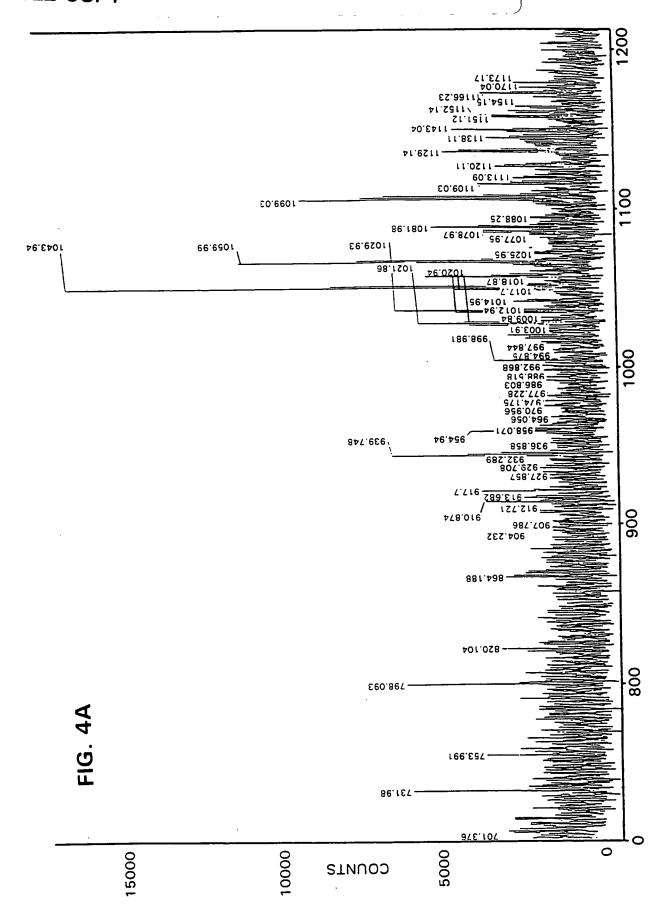
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Appln No.: 10/808,041 Applicant(s): Roman M. Chicz et al.

PROFILING AND CATALOGING EXPRESSED PROTEIN

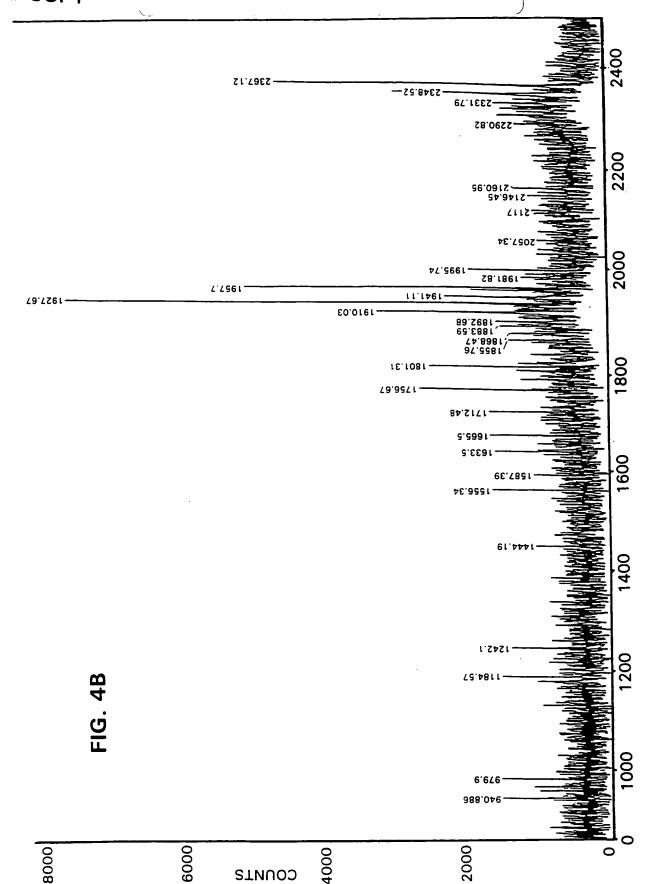
Page 5 of 10

TAGS



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Appln No.: 10/808,041 Page 6 of Applicant(s): Roman M. Chicz et al. PROFILING AND CATALOGING EXPRESSED PROTEIN Page 6 of 10

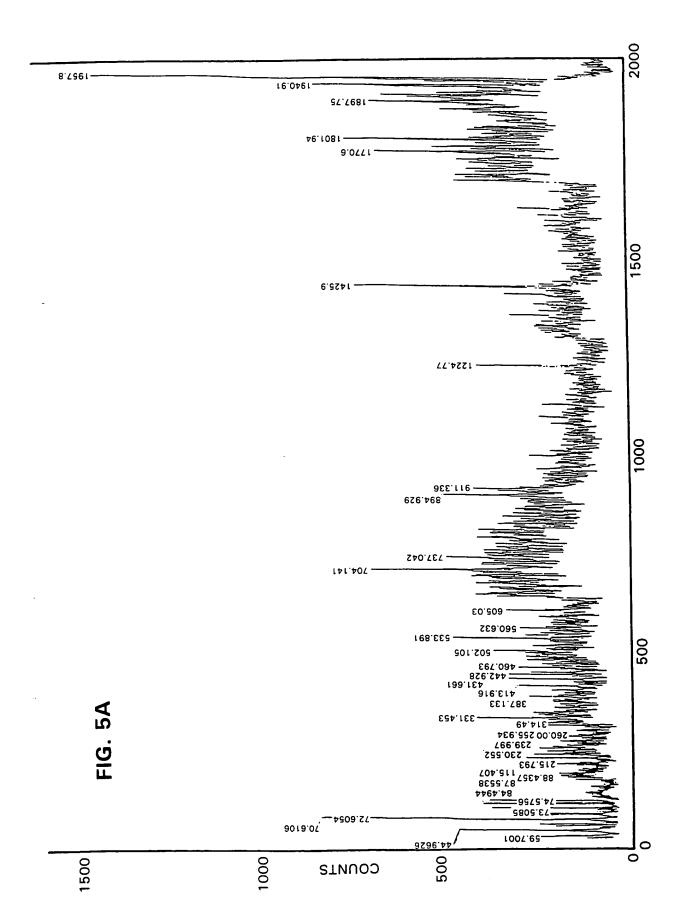


Page 7 of 10

Appln No.: 10/808,041 Applicant(s): Roman M. Chicz et al.

PROFILING AND CATALOGING EXPRESSED PROTEIN

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Page 8 of 10

Appln No.: 10/808,041 Page 8 of Applicant(s): Roman M. Chicz et al.
PROFILING AND CATALOGING EXPRESSED PROTEIN

Sample ID (comment): R4A3F37m1957
Database searched: NCBInr.7.5.97
Molecular weight search (1000-100000 Da) selects 269572 entries.
Species search (HOMO SAPIENS) selects 22771 entries.
Combined molecular weight and species searches select 20933 entries.
Number of sequences passing through parent mass filter: 84015
Ion Types Considered: a b B y n h I

search selects 257 entries

Peptide C terminus Free Acid (O II)

Peptide y N terminus Hydrogen (II)

Cysteines Modified by unmodified

Max. # Missed Cleavages

Peptide Masses are Par(ml)Frag(av)

Search Max. # Mode Unmatched Ions Identity 5

No enzyme Digest Used

											chain
	Protein Name	(D82930) HLA-A26-varient	(U25971) MHC class 1 antigen HLA-A2407	(X82161) HLA-A alpha1 and alpha2 domains	(D32129) HLA-A26	(297370) human leukocyte antigen	(U18930)MHC class 1 antigen HLA-A2	(M160010) HLA-AH class 1 antigen (AA at 30)	(AF017310) MHC class 1 antigen	(AF017309) MHC class 1 antigen	(AF012767) MHC class 1 antigen HLA-A heavy chain
Result Summary	# Unmatched Ions	2	2	7	7	7	7	2	2	2	2
	Sequence	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	0.2948 (Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)	(Y)VDDTQFVRFDSDAASQR(M)
	MH+ Error (Da)	0.2948	0.2948	0.2948	0.2948	0.2948	0.2948	0.2948	0.2948	0.2948	0.2948
	Calculated MH+ (Da)	1956.9052	1956.9052	1956.9052	1956.9052	1956.9052	1956.9052	1956.9052	1956.9052	1956.9052	1956.9052
	Species	HOMO	HOMO	HOMO	HOMO	HOMO	HOMO	HOMO	HOMO	HOMO	HOMO
Parent mass. 1957.2000 (+/-500.0000ppm) Fragment Ions present: [RPJV[NR] KQ]F5DR	Protein MW (Da)	21251.4	40679.8	20921.2	41113.1	38703.4	40895.1	38355.8	10423.4	10430.3	21027.3
	MS-Digest NCBInr.7.597 Index# Accession#	1839795	915219	825673	994765	250934	717123	785055	2394324	2394322	2394009
	MS-Digest Index#	240839	152874	141923	159176	282322	133479	49436	277806	277805	277713
Parent Fragme	Rank	-	-	-	-	-	-	-	-	-	-

qb/R12066/R12066 yf54d10.rl Homo sapiens cDNA clone 26062 5'similar to qb:X00492 cds 1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-3A*0301 alpha (human);

Plus Strand HSPs:

= +2 Identities = 18/18 (100%), Positives=18/18(100%), Frame Score = 90 (41.9 bits), Expect = 0.1e-05, P=9.1e-05

VDDTQFVRFDSDAASQRM -Query:

18

158 VDDTQFVRFDSDAASQRM 211 VDDTQFVRFDSDAASQRM Sbjct:

sapiens

Length = 428

Plus Strand HSPs:

Score = 90 (41.9) bits), Expect = 8.1e-05,P=8.1e-05 Identities = 18/18 (100%) Positives=18/18(100%) Frame = +3

Query: 1 VDDTQFVRFDSDAASQRM 18

VDDTQFVRFDSDAASQRM

Sbjct: 158 VDDTQFVRFDSDAASQRM 203

9764/R59764_yh07c05.rl Homo sapiens cDNA clone 42563 5'similar to gb:X00492_cds1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN,A-3 A* 0301 ALPHA (HUMAN):. qb/R59764/R59764

Length = 461

Plus Strands HSPs:

Score = 90 (41.9 bits), Expect = 8.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame =+1

138 VDDTQFVRFDSDAASQRM 210 Query: 1 VDDTQFVRFDSDAASQRM 18 VDDTQFVRFDSDAASQRM

gb/AA488534/AA488534 ab37f08.rl Stratagene HeLa cell s3 937216 Homo sapiens CDNA clone 843015 5' similar to gb:L06425 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, AW-34(A-10)) A*3402 ALPHA (HUNAN): Length = 478

Plus Strand HSPs:

+3 Score = 90 (41.9 bits), Expect = 8.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame

18 1 VDDTQFVRFDSDAASQRM VDDTQFVRFDSDAASQRM Query:

138 VDDTQFVRFDSDAASQRM 191 Sbjct:

gb/AA548636/AA548636 nj 38d02.sl NCI CGAP AA1 Homo Sapiens cDNA clone 1MAGE 994755 similar to gb:z46633 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-2 a 0201 ALPHA (HUMAN); Length = 539

Plus Strand HSPs:

= +2 Score= 30 (41.9 bits), Expect = 8.1e-05, p = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame

18 VDDTQFVRFDSDAASQRM -Query:

VDDTQFVRFDSDAASQRM 205 VDDTQFVRFDSDAASQRM Sbjct: 152 qb/AA147151/AA147151 zo32d06.rl Stratagene colon (#937204) Homo sapiens cDNA clone 5885875 similar to gb:M64740 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-24 (A-9) A*2402 ALPHA (HUMAN);

Length - 581

Plus Strands HSPs:

Score = 90 (41.9) bits), Expect = 8.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame = +1

1 VDDTQFVRFDSDAASQRM 18 VDDTQFVRFDSDAASQRM Query:

13 VDDTQFVRFDSDAASQRM 66

qb/ H23377/H23377 ym57e02.r1 Homo sapiens cDNA clone 52227 5'similar to qb:x00492 cds1 HIA CLASS I HISTOCOMPATIBILITY ANTIGEN,A-3A A*0301 ALPHA (HUMAN);

Length = 459

Plus Strand HSPs:

Score = 90 (41.9 bits), Expect = 8.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame

7

1 VDDTQEVREDSDAASQRM 18 VDDTQEVREDSDAASQRM Query:

154 VDDTQFVRFDSDAASQRM 207 Sbjct: qb/R13904/R13904 yf62c03.rl Homo sapiens cDNA clone 26801 5'similar to qb:m64742_cds1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-23(A-9) A+2301 (HUMAN);

Length = 459

Plus Strand HSPs:

Score 90 (41.9 bits), Expect = 8.1e-05 P=8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame = +2

18 1 VDDTQFVRFDSDAASQRM VDDTQFVRFDSDAASQRM Query:

Sbjct: 146 VDDTQFVRFDSDAASQRM 199

Appln No.: 10/808,041 Page 10 of Applicant(s): Roman M. Chicz et al. PROFILING AND CATALOGING EXPRESSED PROTEIN Page 10 of 10

